

SEQUENCE LISTING

<110> SHEFFIELD, VAL C.
 ALWARD, WALLACE L.M.
 STONE, EDWIN M.
 NISHIMURA, DARRYL
 PATIL, SHIVA

<120> THERAPEUTICS AND DIAGNOSTICS FOR CONGENITAL HEART DISEASE BASED ON A NOVEL HUMAN TRANSCRIPTION FACTOR

RECFI

0015116

TECH CENTER (600 21 L)

<130> IOWA:042USD1

<140> 09/612,809

<141> 2000-07-10

<160> 20

<170> PatentIn Ver. 2.1

<210> 1

<211> 2284

<212> DNA

<213> Homo sapiens

<400> 1

cgagaaaagg tgacgcgggg cccgggcagg cggccggcgc gcggccccc cccccccgc 60 cctggttatt tggccgcctt cgccggcagc tcagggcaga gtctcctgga aggcgcaggc 120 agtgtggcga gaagggcgcc tgcttgttct ttctttttgt ctgctttccc ccgtttgcgc 180 ctggaagetg egeegegagt teetgeaagg eggtetgeeg eggeegggee eggeettete 240 ccctcgcagc gaccccgct cgcggccgcg cgggccccga ggtagcccga ggcgccggag 300 gagecagece cagegagege egggagagege ggcagegeag eeggaegeac agegeagegg 360 geeggeacea geteggeegg geeggaete ggaeteggeg geeggeegg egeggeeegg 420 cccgagcgag ggtgggggc ggcgggcggc gcggggcggc ggcgagcggg ggcccacacc 480 ctcaaagccg aactaaatcg aaccccaaag caggaaaagc taaaggaacc catcaaggca 540 aaatcgaaac taaaaaaaaa aaatccaatt aaaaaaaacc cctgagaata ttcaccacac 600 caqcqaacag aatatccctc caaaaattca gctcaccagc accagcacga agaaaactct 660 attttcttaa ccgattaatt cagagccacc tccactttgc cttgtctaaa taaacaaacc 720 cgtaaactgt tttatacaga gacagcaaaa tcttggttta ttaaaggaca gtgttactcc 780 agataacacg taagtttett ettgetttte agagacetge ttteecetee teeegtetee 840 cctctcttgc cttcttcctt gcctctcacc tgtaagatat tattttatcc tatgttgaag 900 ggagggggaa agtccccgtt tatgaaagtc gctttctttt tattcatgga cttgttttaa 960 aatgtaaatt gcaacatagt aatttatttt taatttgtag ttggatgtcg tggaccaaac 1020 gccagaaagt gttcccaaaa cctgacgtta aattgcctga aactttaaat tgtgcttttt 1080 ttctcattat aaaaagggaa actgtattaa tcttattcta tcctcttttc tttctttttg 1140 ttgaacatat tcattgtttg tttattaata aattaccatt cagtttgaat gagacctata 1200 tgtctggata ctttaataga gctttaatta ttacgaaaaa agatttcaga gataaaacac 1260 tagaagttac ctattctcca cctaaatctc tgaaaaatgg agaaaccctc tgactagtcc 1320 atgicaaatt tiactaaaag tottittigit tagattiatt ticctgcago atottotgca 1380 aaatgtacta tataqtcagc ttgctttgag gctagtaaaa agatattttt ctaaacagat 1440 tggagttggc atataaacaa atacgttttc tcactaatga cagtccatga ttcggaaatt 1500 ttaagcccat gaatcagccg cggtcttacc acggtgatgc ctgtgtgccg agagatggga 1560 ctgtgcggcc agatatgcac agataaatat ttggcttgtg tattccatat aaaattgcag 1620 tgcatattat acatccctgt gagccagatg ctgaatagat tttttcctat tatttcagtc 1680 ctttataaaa ggaaaaataa accagttttt aaatgtatgt atataattct cccccattta 1740 caatccttca tgtattacat agaaggattg cttttttaaa aatatactgc gggttggaaa 1800 gggatattta atctttgaga aactatttta gaaaatatgt ttgtagaaca attatttttg 1860 aaaaagattt aaagcaataa caagaaggaa ggcgagagga gcagaacatt ttggtctagg 1920 qtqqtttctt tttaaaccat tttttcttqt taatttacaq ttaaacctaq qqqacaatcc 1980 ggattggccc tccccctttt gtaaataacc caggaaatgt aataaattca ttatcttagg 2040 gtgatctgcc ctgccaatca gactttgggg agatggcgat ttgattacag acgttcgggg 2100 gggtgggggg cttgcagttt gttttggaga taatacagtt tcctgctatc tgccgctcct 2160 atctagaggc aacacttaag cagtaattgc tgttgcttgt tgtcaaaatt tgatcattgt 2220 aaaa 2284

<210> 2

<211> 553

<212> PRT

<213> Homo sapiens

<400> 2

Met Gln Ala Arg Tyr Ser Val Ser Ser Pro Asn Ser Leu Gly Val Val 1 5 10 15

Pro Tyr Leu Gly Gly Glu Gln Ser Tyr Tyr Arg Ala Ala Ala Ala Ala 20 25 30

Ala Gly Gly Tyr Thr Ala Met Pro Ala Pro Met Ser Val Tyr Ser 35 40 45

His Pro Ala His Ala Glu Gln Tyr Pro Gly Gly Met Ala Arg Ala Tyr 50 55 60

Gly Pro Tyr Thr Pro Gln Pro Gln Pro Lys Asp Met Val Lys Pro Pro 65 70 75 80

Tyr Ser Tyr Ile Ala Leu Ile Thr Met Ala Ile Gln Asn Ala Pro Asp 85 90 95

Lys Lys Ile Thr Leu Asn Gly Ile Tyr Gln Phe Ile Met Asp Arg Phe 100 105 110

Pro Phe Tyr Arg Asp Asn Lys Gln Gly Trp Gln Asn Ser Ile Arg His
115 120 125

Asn Leu Ser Leu Asn Glu Cys Phe Val Lys Val Pro Arg Asp Asp Lys Lys Pro Gly Lys Gly Ser Tyr Trp Thr Leu Asp Pro Asp Ser Tyr Asn Met Phe Glu Asn Gly Ser Phe Leu Arg Arg Arg Arg Phe Lys Lys Lys Asp Ala Val Lys Asp Lys Glu Glu Lys Asp Arg Leu His Leu Lys Glu Pro Pro Pro Gly Arg Gln Pro Pro Pro Ala Pro Pro Glu Gln Ala Asp Gly Asn Ala Pro Gly Pro Gln Pro Pro Pro Val Arg Ile Gln Asp Ile Lys Thr Glu Asn Gly Thr Cys Pro Ser Pro Pro Gln Pro Leu Ser Pro Ala Ala Ala Leu Gly Ser Gly Ser Ala Ala Ala Val Pro Lys Ile Glu Ser Pro Asp Ser Ser Ser Ser Ser Leu Ser Ser Gly Ser Ser Pro Pro Gly Ser Leu Pro Ser Ala Arg Pro Leu Ser Leu Asp Gly Ala Asp Ser Ala Pro Pro Pro Pro Ala Pro Ser Ala Pro Pro Pro His His Ser Gln Gly Phe Ser Val Asp Asn Ile Met Thr Ser Leu Arg Gly Ser Pro Gln Ser Ala Ala Ala Glu Leu Ser Ser Gly Leu Leu Ala Ser Ala Ala Ala Ser Ser Arg Ala Gly Ile Ala Pro Pro Leu Ala Leu Gly Ala Tyr Ser Pro Gly Gln Ser Ser Leu Tyr Ser Ser Pro Cys Ser Gln Thr Ser Ser Ala Gly Ser Ser Gly Gly Gly Gly Gly Ala Gly Ala Ala

Gly Gly Ala Gly Gly Ala Gly Thr Tyr His Cys Asn Leu Gln Ala Met 385 390 395 400

Ser Leu Tyr Ala Ala Gly Glu Arg Gly Gly His Leu Gln Gly Ala Pro 405 410 415

Gly Gly Ala Gly Gly Ser Ala Val Asp Asn Pro Leu Pro Asp Tyr Ser 420 425 430

Leu Pro Pro Val Thr Ser Ser Ser Ser Ser Ser Leu Ser His Gly Gly
435 440 445

Gly Gly Gly Gly Gly Gly Gln Glu Ala Gly His His Pro Ala 450 455 460

Ala His Gln Gly Arg Leu Thr Ser Trp Tyr Leu Asn Gln Ala Gly Gly
465 470 475 480

Asp Leu Gly His Leu Ala Ser Ala Ala Ala Ala Ala Ala Ala Gly
485 490 495

Tyr Pro Gly Gln Gln Gln Asn Phe His Ser Val Arg Glu Met Phe Glu 500 505 510

Ser Gln Arg Ile Gly Leu Asn Asn Ser Pro Val Asn Gly Asn Ser Ser 515 520 525

Cys Gln Met Ala Phe Pro Ser Ser Gln Ser Leu Tyr Arg Thr Ser Gly 530 535 540

Ala Phe Val Tyr Asp Cys Ser Lys Phe 545 550

<210> 3

<211> 1662

<212> DNA

<213> Homo sapiens

<400> 3

cgcgacgaca agaagccggg caagggcagc tactggacgc tggacccgga ctcctacaac 480 atgttcgaga acggcagctt cctgcggcgg cggcggcgct tcaagaagaa ggacgcggtg 540 aaggacaagg aggagaagga caggctgcac ctcaaggagc cgccccgcc cggccgccag 600 cccccgcccg cgccgccgga gcaggccgac ggcaacgcgc ccggtccgca gccgccgccc 660 gtgcgcatcc aggacatcaa gaccgagaac ggtacgtgcc cctcgccgcc ccagcccctg 720 tecceggeeg eegecetggg eageggeage geegeeggg tgeeeaagat egagageeee 780 gacagcagca gcagcagcct gtccagcggg agcagcccc cgggcagcct gccgtcggcg 840 eggeegetea geetggaegg tgeggattee gegeegeege egeeggege eteegeeeeg 900 ccgccgcacc atagccaggg cttcagcgtg gacaacatca tgacgtcgct gcgggggtcg 960 cegeagageg eggeegega geteagetee ggeettetgg ceteggegge egegteeteg 1020 cgcgcgggga tcgcaccccc gctggcgctc ggcgcctact cgcccggcca gagctccctc 1080 tacagetece eetgeageca gacetecage gegggeaget egggeggegg eggeggegge 1140 gegggggeeg eggggggege gggeegeec gggaectace actgcaacet gcaagecatg 1200 agcctgtacg cggccggcga gcgcggggc cacttgcagg gcgcgcccgg gggcgcgggc 1260 ggctcggccg tggacaaccc cctgcccgac tactctctgc ctccggtcac cagcagcagc 1320 tegtegtece tgagteaegg eggeggegg ggeggeggeg ggggaggeea ggaggeegge 1380 caccaccetg eggeecacca aggeegeete acetegtggt acetgaacca ggegggegga 1440 gacctgggcc acttggcaag cgcggcggcg gcggcggcgg ccgcaggcta cccgggccag 1500 cagcagaact tccactcggt gcgggagatg ttcgagtcac agaggatcgg cttgaacaac 1560 totocagtga acgggaatag tagotgtoaa atggcottoo ottocagooa gtototgtac 1620 cgcacgtccg gagctttcgt ctacgactgt agcaagtttt ga 1662

<210> 4

<211> 106

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 4

Pro Lys Asp Met Val Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Thr
1 5 10 15

Met Ala Ile Gln Asn Ala Pro Asp Lys Lys Ile Thr Leu Asn Gly Ile
20 25 30

Tyr Gln Phe Ile Met Asp Arg Phe Pro Phe Tyr Arg Asp Asn Lys Gln 35 40 45

Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Glu Cys Phe
50 55 60

Val Lys Val Pro Arg Asp Asp Lys Lys Pro Gly Lys Gly Ser Tyr Trp
65 70 75 80

Thr Leu Asp Pro Asp Ser Tyr Asn Met Phe Glu Asn Gly Ser Phe Leu

85 90 95

Arg Arg Arg Arg Phe Lys Lys Lys Asp 100 105

<210> 5

<211> 106

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 5

Pro Lys Asp Leu Val Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Thr 1 5 10 15

Met Ala Ile Gln Asn Ala Pro Glu Lys Lys Ile Thr Leu Asn Gly Ile 20 25 30

Tyr Gln Phe Ile Met Asp Arg Phe Pro Phe Tyr Arg Glu Asn Lys Gln 35 40 45

Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Glu Cys Phe
50 55 60

Val Lys Val Pro Arg Asp Asp Lys Lys Pro Gly Lys Gly Ser Tyr Trp 65 70 75 80

Thr Leu Asp Pro Asp Ser Tyr Asn Met Phe Glu Asn Gly Ser Phe Leu 85 90 95

Arg Arg Arg Arg Phe Lys Lys Asp 100 105

<210> 6

<211> 106

<212> PRT

<213> Artificial Sequence

<220>

<400> 6 Thr Thr Glu Pro Thr Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Ala 10 Met Ala Ile Gln Ser Ser Pro Gly Gln Arg Ala Thr Leu Ser Gly Ile 20 25 30 Tyr Arg Val Ile Met Gly Arg Phe Ala Phe Tyr Arg His Asn Arg Pro 40 Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Glu Cys Phe 55 Val Lys Val Pro Arg Asp Asp Arg Lys Pro Gly Lys Gly Ser Tyr Trp 70 75 Thr Leu Asp Pro Asp Cys His Asp Met Phe Glu His Gly Ser Phe Leu Arg Arg Arg Arg Phe Thr Arg Gln Thr 100 <210> 7 <211> 106 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide Ala Glu Thr Pro Gln Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Ala 10 Met Ala Ile Gln Asp Ala Pro Glu Gln Arg Val Thr Leu Asn Gly Ile 25 Tyr Gln Phe Ile Met Asp Arg Phe Pro Phe Tyr His Asp Asn Arg Gln 35 40 45 Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Asp Cys Phe 50 55

Val Lys Val Pro Arg Glu Lys Gly Arg Pro Gly Lys Gly Ser Tyr Trp

75

80

70

Thr Leu Asp Pro Arg Cys Leu Asp Met Phe Glu Asn Gly Asn Tyr Arg
85 90 95

Arg Arg Lys Arg Lys Pro Lys Pro Gly Pro 100 105

<210> 8

<211> 106

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 8

Pro Leu Gln Arg Gly Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Ala 1 5 10 15

Met Ala Leu Ala His Ala Pro Gly Arg Arg Leu Thr Leu Ala Ala Ile 20 25 30

Tyr Arg Phe Ile Thr Glu Arg Phe Ala Phe Tyr Arg Asp Ser Pro Arg
35 40 45

Lys Trp Gln Asn Ser Ile Arg His Asn Leu Thr Leu Asn Asp Cys Phe 50 55 60

Val Lys Val Pro Arg Glu Pro Gly Asn Pro Gly Lys Gly Asn Tyr Trp
65 70 75 80

Thr Leu Asp Pro Ala Ala Ala Asp Met Phe Asp Asn Gly Ser Phe Leu 85 90 95

Pro Arg Arg Lys Arg Phe Lys Arg Ala Glu 100 105

<210> 9

<211> 106

<212> PRT

<213> Artificial Sequence

<220>

1	•)
			•			,									
)> 9 Leu	Gln	Arg	Gly 5	Lys	Pro	Pro	Tyr	Ser 10	Tyr	Ile	Ala	Leu	Ile 15	A]
Met	Ala	Ile	Ala 20	His	Ala	Pro	Glu	Arg 25	Arg	Leu	Thr	Leu	Gly 30	Gly	I
Tyr	Lys	Phe 35	Ile	Thr	Glu	Arg	Phe 40	Pro	Phe	Tyr	Arg	Asp 45	Asn	Pro	Ŀ
Lys	Trp 50	Gln	Asn	Ser	Ile	Arg 55	His	Asn	Leu	Thr	Leu 60	Asn	Asp	Cys	P
Leu 65	Lys	Ile	Pro	Arg	Glu 70	Ala	Gly	Arg	Pro	Gly 75	Lys	Gly	Asn	Tyr	Т
Ala	Leu	Asp	Pro	Asn 85	Ala	Glu	Asp	Met	Phe 90	Glu	Ser	Gly	Ser	Phe 95	L
Arg	Arg	Arg	Lys 100	Arg	Phe	Lys	Arg	Ser 105	Asp						
<21 <21	0 > 1 1 > 1 2 > P 3 > A	.06	icia	l Se	quen	ce									
<22 <22	3 > E	escr Pepti		on o	f Ar	tifi	cial	Seq	uenc	e: :	Synt]	heti	С		
<40	0 > 1	. 0													
	Arg	g Gln	Pro	Ala 5		Pro	Pro	Ser	Ser 10		Ile	Ala	Leu	Ile 15	
Met	Ala	a Ile	Leu 20		Ser	Pro	His	Lys 25	Arg	Leu	Thr	Leu	Ser 30]
Cys	Ala	a Phe 35		Ser	Asp	Arg	Phe		Tyr	Tyr	Arg	Arg		Glu	F
Gly	Trp	o Gln	Asn	. Ser	Ile	Arg 55		Asn	Leu	Ser	Leu 60		Asp	Cys	. !
		s Ile													

Ser Leu Asp Pro Ala Ser Gln Asp Met Phe Asp Asn Gly Ser Phe Leu 90 95 85 Arg Arg Arg Lys Arg Phe Gln Arg Asn Gln 100 105 <210> 11 <211> 106 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 11 Arg Thr Arg Leu Val Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Thr Met Ala Ile Leu Gln Ser Pro Lys Lys Arg Leu Thr Leu Ser Glu Ile 20 25 30 Cys Glu Phe Ile Ser Gly Arg Phe Pro Tyr Tyr Arg Glu Lys Phe Pro 35 40 Ala Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Asp Cys Phe 55 60 50 Val Lys Ile Pro Arg Glu Pro Gly Asn Pro Gly Lys Gly Asn Tyr Trp 65 70 75 Thr Leu Asp Pro Glu Ser Ala Asp Met Phe Asp Asn Gly Ser Phe Leu 85 90 95 Arg Arg Arg Lys Arg Phe Lys Arg Gln Pro 100 105 <210> 12 <211> 106 <212> PRT <213> Artificial Sequence

<400> 12 Arg Ser Pro Leu Val Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Thr 10 5 Met Ala Ile Leu Gln Ser Pro Lys Lys Arg Leu Thr Leu Ser Glu Ile 25 20 Cys Glu Phe Ile Ser Gly Arg Phe Pro Tyr Tyr Arg Glu Lys Phe Pro 35 40 Ala Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Asp Cys Phe 55 Val Lys Ile Pro Arg Glu Pro Gly Asn Pro Gly Lys Gly Asn Tyr Trp 70 75 Thr Leu Asp Pro Glu Ser Ala Asp Met Phe Asp Asn Gly Ser Phe Leu 85 Arg Arg Lys Arg Arg Phe Lys Arg Gln Pro 100 <210> 13 <211> 106 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 13 Ile Arg Arg Pro Glu Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Val 10 Met Ala Ile Gln Ser Ser Pro Thr Lys Arg Leu Thr Leu Ser Glu Ile 25 Tyr Gln Phe Leu Gln Ser Arg Phe Pro Phe Phe Arg Gly Ser Tyr Gln 35 40 45 Gly Trp Lys Asn Ser Val Arg His Asn Leu Ser Leu Asn Glu Cys Phe 50 55 Ile Lys Leu Pro Lys Gly Leu Gly Arg Pro Gly Lys Gly His Tyr Trp

65

70

Thr Ile Asp Pro Ala Ser Glu Phe Met Phe Glu Asn Gly Ser Phe Arg 85 90 95 Arg Arg Arg Gly Phe Arg Arg Lys Cys 100 <210> 14 <211> 106 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide Leu Arg Arg Pro Glu Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Val 15 10 Met Ala Ile Gln Ser Ser Pro Ser Lys Arg Leu Thr Leu Ser Glu Ile 30 25 20 Tyr Gln Phe Leu Gln Ala Arg Phe Pro Phe Phe Arg Gly Ala Tyr Gln 40 45 35 Gly Trp Lys Asn Ser Val Arg His Asn Leu Ser Leu Asn Glu Cys Phe 55 60 50 Ile Lys Leu Pro Lys Gly Leu Gly Arg Pro Gly Lys Gly His Tyr Trp 65 70 75 Thr Ile Asp Pro Ala Ser Glu Phe Met Phe Glu Asn Gly Ser Phe Arg 90 95 85 Arg Arg Arg Gly Phe Arg Arg Lys Cys 100 105 <210> 15 <211> 106 <212> PRT <213> Artificial Sequence

<220>

<400> 15 Asn Gly Lys Tyr Glu Lys Pro Pro Phe Ser Tyr Asn Ala Leu Ile Met 10 Met Ala Ile Arg Gln Ser Pro Glu Lys Arg Leu Thr Leu Asn Gly Ile 20 25 Tyr Glu Phe Ile Met Lys Asn Phe Pro Tyr Tyr Arg Glu Asn Lys Gln 40 35 Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Lys Cys Phe 55 Val Lys Val Pro Arg His Tyr Asp Asp Pro Gly Lys Gly Asn Tyr Trp 70 Met Leu Asp Pro Ser Ser Tyr Asp Asp Val Ile Gly Gly Thr Thr Gly 85 90 Lys Leu Arg Arg Ser Thr Thr Ser Pro 100 105 <210> 16 <211> 106 <212> PRT <213> Artificial Sequence <400> 16 Asn Gly Lys Tyr Glu Lys Pro Pro Phe Ser Tyr Asn Ala Leu Ile Met 10 Met Ala Met Arg Gln Ser Pro Glu Lys Arg Leu Thr Leu Asn Gly Ile 20 25 Tyr Glu Phe Ile Met Lys Asn Phe Pro Tyr Tyr Arg Glu Asn Lys Gln Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Lys Cys Phe 55 Val Lys Val Pro Arg His Tyr Asp Asp Pro Gly Lys Gly Asn Tyr Trp 65 70 75 Met Leu Asp Pro Ser Ser Tyr Asp Asp Val Ile Gly Gly Thr Thr Gly

90

Lys Leu Arg Arg Ser Thr Thr Ser Pro Ala

100 105

<210> 17 <211> 106 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 17 Gly Lys Tyr Glu Lys Pro Pro Pro Phe Ser Tyr Asn Ala Leu Ile Met Met Ala Ile Arg Gln Ser Pro Glu Lys Arg Leu Thr Leu Asn Gly Ile 25 20 Tyr Glu Phe Ile Met Lys Asn Phe Pro Tyr Tyr Arg Glu Asn Lys Gln 35 40 Gly Trp His Asn Ser Ile Arg Asp Asn Leu Ser Leu Asn Lys Cys Phe 60 55 50 Val Lys Val Pro Arg His Tyr Asp Asp Pro Gly Lys Gly Asn Tyr Trp 80 75 70 65 Met Leu Asp Pro Ser Ser Asp Asp Val Phe Ile Gly Gly Thr Thr Gly 95 90 85 Lys Leu Arg Arg Arg Ser Thr Thr Ser Arg 105 100 <210> 18 <211> 76 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 18

Leu Met Lys Leu Val Arg Pro Pro Tyr Ser Tyr Ser Ala Leu Ile Ala

5

1

10

Met Ala Ile His Gly Ala Pro Asp Lys Arg Leu Thr Leu Ser Gln Ile 25 20 Tyr Gln Tyr Val Ala Asp Asn Phe Pro Phe Tyr Asn Lys Ser Lys Ala 35 40 Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Asp Cys Phe 50 55 Lys Lys Val Pro Arg Asp Glu Asp Asp Pro Gly Lys 70 65 <210> 19 <211> 106 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 19 Thr Asn Pro His Val Lys Pro Pro Tyr Ser Tyr Ala Thr Leu Ile Cys 10 Met Ala Met Gln Ala Ser Lys Ala Thr Lys Ile Thr Leu Ser Ala Ile 20 25 30 Tyr Lys Trp Ile Thr Asp Asn Phe Cys Tyr Phe Arg His Ala Asp Pro 35 40 45 Thr Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Lys Cys Phe 55 Ile Lys Val Pro Arg Glu Lys Asp Glu Pro Gly Lys Gly Phe Trp Arg Ile Asp Pro Gln Tyr Ala Glu Arg Leu Leu Ser Gly Ala Phe Lys 85 90 Lys Arg Arg Leu Pro Phe Val His Ile His 100 <210> 20 <211> 98

<212> PRT

<213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 20 Trp Gly Asn Leu Ser Tyr Ala Asp Leu Ile Thr Lys Ala Ile Glu Ser 10 Ser Ala Glu Lys Arg Leu Thr Leu Ser Gln Ile Tyr Glu Trp Met Val 20 25 Lys Ser Val Pro Tyr Phe Lys Asp Lys Gly Asp Ser Asn Ser Ser Ala 40 Gly Trp Gln Lys Ser Ile Arg His Asn Leu Ser Leu His Ser Lys Phe 55 Ile Arg Val Gln Asn Glu Gly Thr Gly Lys Ser Ser Trp Trp Met Leu 75 65 70 80 Asn Pro Glu Gly Gly Lys Ser Gly Lys Ser Pro Arg Arg Ala Ala Ser 85 90 95

Met Asp